

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/581,810
Source: IFW
Date Processed by STIC: 3/14/07

ENTERED



IFWO

RAW SEQUENCE LISTING

DATE: 03/14/2007

PATENT APPLICATION: US/10/581,810

TIME: 14:58:16

Input Set : N:\efs\03_14_07\10581810_efs\HERR13001APCSEQLIST.TXT

Output Set: N:\CRF4\03142007\J581810.raw

```

4 <110> APPLICANT: Ortega Mora, Luis MIGuel
5      Fernandez Garcia, Aurora
8 <120> TITLE OF INVENTION: Use of gene NcSAG4 for the diagnosis and
9      prevention of neosporosis and as a marker for analysis of
10     the pathogenesis
12 <130> FILE REFERENCE: HERR13.001APC
14 <140> CURRENT APPLICATION NUMBER: US 10/581,810
15 <141> CURRENT FILING DATE: 2006-06-02
17 <150> PRIOR APPLICATION NUMBER: PCT/ES2004/000529
18 <151> PRIOR FILING DATE: 2004-11-26
20 <150> PRIOR APPLICATION NUMBER: ES P200302869
21 <151> PRIOR FILING DATE: 2003-12-04
23 <160> NUMBER OF SEQ ID NOS: 15
25 <170> SOFTWARE: FastSEQ for Windows Version 4.0
27 <210> SEQ ID NO: 1
28 <211> LENGTH: 23
29 <212> TYPE: DNA
30 <213> ORGANISM: Neospora caninum
32 <220> FEATURE:
33 <221> NAME/KEY: modified_base
34 <222> LOCATION: 6
35 <223> OTHER INFORMATION: I
37 <400> SEQUENCE: 1
W--> 38 tggacntayg ayttyaaraa rgc                23
40 <210> SEQ ID NO: 2
41 <211> LENGTH: 23
42 <212> TYPE: DNA
43 <213> ORGANISM: Neospora caninum
45 <220> FEATURE:
46 <221> NAME/KEY: modified_base
47 <222> LOCATION: 12, 15, 18, 21
48 <223> OTHER INFORMATION: I
50 <400> SEQUENCE: 2
W--> 51 aaraargara tnatnacncc ngg                23
53 <210> SEQ ID NO: 3
54 <211> LENGTH: 23
55 <212> TYPE: DNA
56 <213> ORGANISM: Neospora caninum
58 <220> FEATURE:
59 <221> NAME/KEY: modified_base
60 <222> LOCATION: 3
61 <223> OTHER INFORMATION: I
63 <400> SEQUENCE: 3

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W--> 64 acnggytcrt cyttcartg rtc                23
      66 <210> SEQ ID NO: 4
      67 <211> LENGTH: 24
      68 <212> TYPE: DNA
      69 <213> ORGANISM: Neospora caninum
      71 <220> FEATURE:
      72 <221> NAME/KEY: modified_base
      73 <222> LOCATION: 22
      74 <223> OTHER INFORMATION: I
      76 <220> FEATURE:
      77 <221> NAME/KEY: misc_feature
      78 <222> LOCATION: 7
      79 <223> OTHER INFORMATION: n = A,T,C or G
      81 <400> SEQUENCE: 4
W--> 82 rtcyttnacy ttraarcara angg                24
      84 <210> SEQ ID NO: 5
      85 <211> LENGTH: 20
      86 <212> TYPE: DNA
      87 <213> ORGANISM: Neospora caninum
      89 <400> SEQUENCE: 5
      90 ccgacgaagc cctgagaact                    20
      92 <210> SEQ ID NO: 6
      93 <211> LENGTH: 19
      94 <212> TYPE: DNA
      95 <213> ORGANISM: Neospora caninum
      97 <400> SEQUENCE: 6
      98 tgtcgctgt tgggttgta                      19
     100 <210> SEQ ID NO: 7
     101 <211> LENGTH: 24
     102 <212> TYPE: DNA
     103 <213> ORGANISM: Neospora caninum
     105 <400> SEQUENCE: 7
     106 gaaacaagaa aagagactat ctca                24
     108 <210> SEQ ID NO: 8
     109 <211> LENGTH: 20
     110 <212> TYPE: DNA
     111 <213> ORGANISM: Neospora caninum
     113 <400> SEQUENCE: 8
     114 ccaggtgaga gtgtttcgat                    20
     116 <210> SEQ ID NO: 9
     117 <211> LENGTH: 601
     118 <212> TYPE: DNA
     119 <213> ORGANISM: Neospora caninum
     121 <220> FEATURE:
     122 <221> NAME/KEY: CDS
     123 <222> LOCATION: (48)...(568)
     125 <400> SEQUENCE: 9
     126 ggcaacacgt cgcagcgtac tctcatcttt ttcgtggggt tgcagcc atg gag aaa    56
     127                                     Met Glu Lys

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128                                     1
130 agc gcc ttc ttt ccc agg gtg gtg ctc tgt ttc gtc gta gtt ttg tcc 104
131 Ser Ala Phe Phe Pro Arg Val Val Leu Cys Phe Val Val Val Leu Ser
132      5                      10                      15
134 gcg tgc tcg gcg tgg cga gtg gaa ggg aag aac tgg tcg tac gat ttc 152
135 Ala Cys Ser Ala Trp Arg Val Glu Gly Lys Asn Trp Ser Tyr Asp Phe
136 20                      25                      30                      35
138 aag aag ccg ctg gac agc gat gaa aca aga aaa gag act atc tca cca 200
139 Lys Lys Pro Leu Asp Ser Asp Glu Thr Arg Lys Glu Thr Ile Ser Pro
140                      40                      45                      50
142 ggt gag agt gtt tcg ata caa aat tct ggg agc att acg ctg gcg tac 248
143 Gly Glu Ser Val Ser Ile Gln Asn Ser Gly Ser Ile Thr Leu Ala Tyr
144                      55                      60                      65
146 aac cca aca ggc gac aca caa gtt ctc agg gct tcg tcg gga gac agc 296
147 Asn Pro Thr Gly Asp Thr Gln Val Leu Arg Ala Ser Ser Gly Asp Ser
148      70                      75                      80
150 tgc agg gat gag cca atc gaa ctt gcg act tta ttc cca gca gcc acg 344
151 Cys Arg Asp Glu Pro Ile Glu Leu Ala Thr Leu Phe Pro Ala Ala Thr
152      85                      90                      95
154 ccg gcg ccc acg tgg atg caa act ggt agc acg aga acc tta gcg ttt 392
155 Pro Ala Pro Thr Trp Met Gln Thr Gly Ser Thr Arg Thr Leu Ala Phe
156 100                      105                      110                      115
158 cct acc aac gca gta ccc gcg aag cag acc acg ccg ttc tgt ttt aaa 440
159 Pro Thr Asn Ala Val Pro Ala Lys Gln Thr Thr Pro Phe Cys Phe Lys
160                      120                      125                      130
162 gtc acg gat acg cag aag aac aaa act ctg aca gcg ata atc aag gtc 488
163 Val Thr Asp Thr Gln Lys Asn Lys Thr Leu Thr Ala Ile Ile Lys Val
164                      135                      140                      145
166 gcc ggt gcc caa ggc ttg tct gct gct ctg ggg gtc tcc att gga ata 536
167 Ala Gly Ala Gln Gly Leu Ser Ala Ala Leu Gly Val Ser Ile Gly Ile
168                      150                      155                      160
170 cca gct ctt gct ttt gca ctg agt tcg ata ta agggcatgca aacgaataaa 588
171 Pro Ala Leu Ala Phe Ala Leu Ser Ser Ile
172      165                      170
174 tgaggcgact gat 601
176 <210> SEQ ID NO: 10
177 <211> LENGTH: 173
178 <212> TYPE: PRT
179 <213> ORGANISM: Neospora caninum
181 <400> SEQUENCE: 10
182 Met Glu Lys Ser Ala Phe Phe Pro Arg Val Val Leu Cys Phe Val Val
183 1                      5                      10                      15
184 Val Leu Ser Ala Cys Ser Ala Trp Arg Val Glu Gly Lys Asn Trp Ser
185      20                      25                      30
186 Tyr Asp Phe Lys Lys Pro Leu Asp Ser Asp Glu Thr Arg Lys Glu Thr
187      35                      40                      45
188 Ile Ser Pro Gly Glu Ser Val Ser Ile Gln Asn Ser Gly Ser Ile Thr
189      50                      55                      60
190 Leu Ala Tyr Asn Pro Thr Gly Asp Thr Gln Val Leu Arg Ala Ser Ser

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```

191 65          70          75          80
192 Gly Asp Ser Cys Arg Asp Glu Pro Ile Glu Leu Ala Thr Leu Phe Pro
193          85          90          95
194 Ala Ala Thr Pro Ala Pro Thr Trp Met Gln Thr Gly Ser Thr Arg Thr
195          100          105          110
196 Leu Ala Phe Pro Thr Asn Ala Val Pro Ala Lys Gln Thr Thr Pro Phe
197          115          120          125
198 Cys Phe Lys Val Thr Asp Thr Gln Lys Asn Lys Thr Leu Thr Ala Ile
199          130          135          140
200 Ile Lys Val Ala Gly Ala Gln Gly Leu Ser Ala Ala Leu Gly Val Ser
201 145          150          155          160
202 Ile Gly Ile Pro Ala Leu Ala Phe Ala Leu Ser Ser Ile
203          165          170
206 <210> SEQ ID NO: 11
207 <211> LENGTH: 26
208 <212> TYPE: DNA
209 <213> ORGANISM: Neospora caninum
211 <400> SEQUENCE: 11
212 gccaggatcc atggagaaaa ggcgcct
214 <210> SEQ ID NO: 12
215 <211> LENGTH: 31
216 <212> TYPE: DNA
217 <213> ORGANISM: Neospora caninum
219 <400> SEQUENCE: 12
220 ttcgaattcc ttatatcgaa ctcagtgcaa a
222 <210> SEQ ID NO: 13
223 <211> LENGTH: 28
224 <212> TYPE: DNA
225 <213> ORGANISM: Neospora caninum
227 <400> SEQUENCE: 13
228 ttatggatcc ggaagaactg gtcgtacg
230 <210> SEQ ID NO: 14
231 <211> LENGTH: 30
232 <212> TYPE: DNA
233 <213> ORGANISM: Neospora caninum
235 <400> SEQUENCE: 14
236 tttgaattcc ttaggcgacc ttgattatcg
238 <210> SEQ ID NO: 15
239 <211> LENGTH: 155
240 <212> TYPE: PRT
241 <213> ORGANISM: Neospora caninum
243 <400> SEQUENCE: 15
244 Met Arg Gly Ser His His His His His Gly Met Ala Ser Met Thr
245 1          5          10          15
246 Gly Gly Gln Gln Met Gly Arg Asp Leu Tyr Asp Asp Asp Asp Lys Asp
247          20          25          30
248 Arg Trp Ile Arg Lys Asn Trp Ser Tyr Asp Phe Lys Lys Pro Leu Asp
249          35          40          45
250 Ser Asp Glu Thr Arg Lys Glu Thr Ile Ser Pro Gly Glu Ser Val Ser

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251	50	55	60
252	Ile Gln Asn Ser Gly Ser	Ile Thr Leu Ala Tyr Asn Pro Thr Gly Asp	
253	65	70	75 80
254	Thr Gln Val Leu Arg Ala Ser Ser Gly Asp Ser Cys Arg Asp Glu Pro		
255		85	90 95
256	Ile Glu Leu Ala Thr Leu Phe Pro Ala Ala Thr Pro Ala Pro Thr Trp		
257		100	105 110
258	Met Gln Thr Gly Ser Thr Arg Thr Leu Ala Pro Thr Asn Ala Val Pro		
259		115	120 125
260	Ala Lys Gln Thr Thr Pro Phe Cys Phe Lys Val Thr Asp Thr Gln Lys		
261		130	135 140
262	Asn Lys Thr Leu Thr Ala Ile Ile Lys Val Ala		
263	145	150	155

RAW SEQUENCE LISTING ERROR SUMMARY
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 6 ✓
Seq#:2; N Pos. 12, 15, 18, 21 ✓
Seq#:3; N Pos. 3 ✓
Seq#:4; N Pos. 7, 22 ✓

VERIFICATION SUMMARY

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Input Set : N:\efs\03_14_07\10581810_efs\HERR13001APCSEQLIST.TXT

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L:38 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
L:51 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0
L:64 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0
L:82 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0